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         May 05
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         May 15
                 Supporter information for ENCOMPPAT and ENCOMPLIT updated
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         May 19
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NEWS 21 Jun 06 Simultaneous left and right truncation added to CBNB
NEWS 22 Jun 06 PASCAL enhanced with additional data
NEWS 23 Jun 20 2003 edition of the FSTA Thesaurus is now available
NEWS 24 Jun 25 HSDB has been reloaded
NEWS 25 Jul 16 Data from 1960-1976 added to RDISCLOSURE
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NEWS 27 Jul 21 Polymer class term count added to REGISTRY
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                 Right Truncation available
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                New pricing for EUROPATFULL and PCTFULL effective
                 August 1, 2003
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        AUG 13
                 Field Availability (/FA) field enhanced in BEILSTEIN
NEWS 31
        AUG 15
                PATDPAFULL: one FREE connect hour, per account, in
                 September 2003
NEWS 32
        AUG 15
                PCTGEN: one FREE connect hour, per account, in
                 September 2003
        AUG 15 RDISCLOSURE: one FREE connect hour, per account, in
NEWS 33
                 September 2003
NEWS 34
        AUG 15 TEMA: one FREE connect hour, per account, in
                 September 2003
NEWS 35 AUG 18 Data available for download as a PDF in RDISCLOSURE
NEWS 36 AUG 18 Simultaneous left and right truncation added to PASCAL
NEWS 37 AUG 18 FROSTI and KOSMET enhanced with Simultaneous Left and Right
                 Truncation
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NEWS 38 AUG 18 Simultaneous left and right truncation added to ANABSTR

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L3

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AN 1998:59202 AGRICOLA

DN IND21236609

TI The role of UDP-glucose epimerase in carbohydrate metabolism of Arabidopsis.

- AU Dormann, P.; Benning, C.
- AV DNAL (OK710.P68)
- The Plant journal : for cell and molecular biology, Mar 1998. Vol. 13, No. 5. p. 641-652
 Publisher: Oxford : Blackwell Sciences Ltd.

ISSN: 0960-7412

- NTE Includes references
- CY England; United Kingdom
- DT Article
- FS Non-U.S. Imprint other than FAO
- LA English
- L3 ANSWER 2 OF 9 BIOSIS COPYRIGHT 2003 BIOLOGICAL ABSTRACTS INC. on STN
- AN 2000:113541 BIOSIS
- DN PREV200000113541
- ${\tt TI}$ Multiple copies of MRG19 suppress transcription of the GAL1 promoter in a GAL80-dependent manner in Saccharomyces cerevisiae.
- AU Kabir, M. A.; Khanday, F. A.; Mehta, D. V.; Bhat, P. J. (1)
- CS (1) Laboratory of Molecular Genetics, Biotechnology Center, Indian Institute of Technology, Powai Mumbai, 400 076 India
- SO Molecular and General Genetics, (Jan., 2000) Vol. 262, No. 6, pp. 1113-1122.
 ISSN: 0026-8925.
- DT Article
- LA English
- SL English
- L3 ANSWER 3 OF 9 BIOSIS COPYRIGHT 2003 BIOLOGICAL ABSTRACTS INC. on STN
- AN 1997:380459 BIOSIS
- DN PREV199799679662
- TI The substrate inhibition of UDP-D-galactose 4-epimerase as possible source of galactose toxicity for higher plants.
- AU Prosselkov, P. V. (1); Gross, W.; Igamberdiev, A. U. (1); Schnarrenberger, C.
- CS (1) Dep. Plant Physiol. Biochem., Biol. Fac., Voronezh State Univ., Voronezh Russia
- SO Plant Physiology (Rockville), (1997) Vol. 114, No. 3 SUPPL., pp. 35.
 Meeting Info.: PLANT BIOLOGY '97: 1997 Annual Meetings of the American
 Society of Plant Physiologists and the Canadian Society of Plant
 Physiologists, Japanese Society of Plant Physiologists and the Australian
 Society of Plant Physiologists Vancouver, British Columbia, Canada August
 2-6, 1997
 ISSN: 0032-0889.
- DT Conference; Abstract
- LA English
- L3 ANSWER 4 OF 9 BIOSIS COPYRIGHT 2003 BIOLOGICAL ABSTRACTS INC. on STN
- AN 1987:193221 BIOSIS
- DN BA83:101345
- TI GALACTOSE INHIBITS THE CONVERSION OF 1 AMINOCYCLOPROPANE-1-CARBOXYLIC ACID TO ETHYLENE IN AGED TOBACCO LEAF DISCS.
- AU PHILOSOPH-HADAS S; AHARONI N
- CS DEP. FRUIT VEGETABLE STORAGE, AGRIC. RES. ORGANIZATION, VOLCANI CENT., BET DAGAN 50250, ISRAEL.
- SO PLANT PHYSIOL (BETHESDA), (1987) 83 (1), 8-11. CODEN: PLPHAY. ISSN: 0032-0889.
- FS BA; OLD
- LA English
- L3 ANSWER 5 OF 9 CAPLUS COPYRIGHT 2003 ACS on STN
- AN 2000:133844 CAPLUS
- DN 132:178178
- ${\tt TI}$ Galactose utilization as a positive selection marker in the transformation of ${\tt plant}$ cells

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Jorsboe, Morten; Brunstedt, Janne; Jorgensen, Kirsten
TM
PΑ
     Danisco A/S, Den.
SO
     PCT Int. Appl., 86 pp.
     CODEN: PIXXD2
DT
     Patent
LA
     English
FAN.CNT 1
                    KIND DATE
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                                          APPLICATION NO. DATE
     WO 2000009705 A2 20000224
WO 2000009705 A3 20000615
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                                                            19990811
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     WO 1999-IB1465
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                            19990811
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     ANSWER 6 OF 9 CAPLUS COPYRIGHT 2003 ACS on STN
AN
     1981:544921 CAPLUS
DN
     95:144921
TI
     The effect of galactose on the growth of Lemna
     DeKock, P. C.; Cheshire, M. V.; Mundie, C. M.; Inkson, R. H. E.
ΑU
CS
     Dep. Plant Physiol., Macaulay Inst. Soil Res., Craigiebuckler/Aberdeen,
     AB9 2QF, UK
SO
     New Phytologist (1979), 82(3), 679-85
     CODEN: NEPHAV; ISSN: 0028-646X
DТ
     Journal
     English
LA
L3
     ANSWER 7 OF 9 CAPLUS COPYRIGHT 2003 ACS on STN
AN
     1965:404979 CAPLUS
DN
     63:4979
OREF 63:931f-g
    Some aspects of sugar nutrition of excised embryos of Lupinus luteus and
TI
     Brassica oleracea
ΑU
    Hoffmanowa, A.
CS
    Univ. Poznan
SO
    Acta Soc. Botan. Polon. (1964), 33(1), 193-210
DT
    Journal
LA
    English
L3
    ANSWER 8 OF 9 CAPLUS COPYRIGHT 2003 ACS on STN
AN
    1917:12025 CAPLUS
DN
    11:12025
OREF 11:2483e-h
    The toxicity of galactose and mannose for green plants and the
    antagonistic action of other sugars towards these
ΑU
    American Journal of Botany (1917), 4, 430-7
    CODEN: AJBOAA: ISSN: 0002-9122
DT
    Journal
```

LA Unavailable

L3 ANSWER 9 OF 9 CAPLUS COPYRIGHT 2003 ACS on STN

AN 1916:6237 CAPLUS

DN 10:6237

OREF 10:1209q h

TI Toxicity of galactose for certain higher plants

AU Knudson, L.

SO Ann. Missouri Bot. Gardens (1915), 2, 659-66

DT Journal

LA Unavailable

=> FIL STNGUIDE

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L3 ANSWER 9 OF 9 CAPLUS COPYRIGHT 2003 ACS on STN

AB Working with a nutrient culture soln., galactose proved harmful to vetch (Vicia villosa) and peas (Pisum sativum) but the other sugars, glucose, lactose, raffinose, sucrose and maltose were beneficial when used in concn. of 2%. Galactose was harmful when used in concn. of 1% and more; below 1% it had no effect.

=> s 12 and bacteria

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=> s 12 and bacteria L4 1 L2 AND BACTERIA

=> d 14 1 ab

- L4 ANSWER 1 OF 1 AGRICOLA Compiled and distributed by the National Agricultural Library of the Department of Agriculture of the United States of America. It contains copyrighted materials. All rights reserved. (2003) on STN
- Uridine 5'-diphospho-glucose-4-epimerase (UDP-Glc epimerase) catalyses the reversible epimerization of UDP-galactose and UDP-glucose. In contrast to bacteria and yeast, expression of the UDP-Glc epimerase gene in Arabidopsis was found not to be induced by galactose. To elucidate the metabolic role of this enzyme, transgenic Arabidopsis plants expressing the respective cDNA in sense or antisense orientation were constructed, leading to a range of plant lines with different UDP-Glc epimerase activities. No alterations in morphology were observed and the relative amounts of different galactose-containing compounds were not affected if the plants were raised on soil. However, on agar plates in the presence of galactose, the growth of different lines was increasingly repressed with decreasing enzyme activity, and an increase in the UDP-Gal content was observed in parallel, whereas the UDP-Glc content was nearly constant. The amount of galactose in the cell wall was increased in plants with low UDP-Glc epimerase activity grown on galactose, whereas the cellulose content in the leaves was not altered. Furthermore, starch determined at different times of the day was highly abundant in plants with low UDP-Glc epimerase activity in the presence of galactose. It is proposed that low endogenous UDP-Glc epimerase activity is responsible for the galactose toxicity of the wild-type. Possible mechanisms by which the starch content might be modulated are discussed.

=> d 14 1

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AN 1998:59202 AGRICOLA

DN IND21236609

TI The role of UDP-glucose epimerase in carbohydrate metabolism of Arabidopsis.

AU Dormann, P.; Benning, C.

AV DNAL (QK710.P68)

SO The Plant journal : for cell and molecular biology, Mar 1998. Vol. 13, No. 5. p. 641-652

Publisher: Oxford: Blackwell Sciences Ltd.

ISSN: 0960-7412

NTE Includes references

CY England; United Kingdom

DT Article

FS Non-U.S. Imprint other than FAO

LA English

=> s ll and bacteria

L5 3 L1 AND BACTERIA

=> d l5 1-3 ab

L5 ANSWER 1 OF 3 AGRICOLA Compiled and distributed by the National

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Uridine 5'-diphospho-glucose-4-epimerase (UDP-Glc epimerase) catalyses the AB reversible epimerization of UDP-galactose and UDP-glucose. In contrast to bacteria and yeast, expression of the UDP-Glc epimerase gene in Arabidopsis was found not to be induced by galactose. To elucidate the metabolic role of this enzyme, transgenic Arabidopsis plants expressing the respective cDNA in sense or antisense orientation were constructed, leading to a range of plant lines with different UDP-Glc epimerase activities. No alterations in morphology were observed and the relative amounts of different galactose-containing compounds were not affected if the plants were raised on soil. However, on agar plates in the presence of galactose, the growth of different lines was increasingly repressed with decreasing enzyme activity, and an increase in the UDP-Gal content was observed in parallel, whereas the UDP-Glc content was nearly constant. The amount of galactose in the cell wall was increased in plants with low UDP-Glc epimerase activity grown on galactose, whereas the cellulose content in the leaves was not altered. Furthermore, starch determined at different times of the day was highly abundant in plants with low UDP-Glc epimerase activity in the presence of galactose. It is proposed that low endogenous UDP-Glc epimerase activity is responsible for the galactose toxicity of the wild-type. Possible mechanisms by which the starch content might be modulated are discussed.

L5 ANSWER 2 OF 3 BIOSIS COPYRIGHT 2003 BIOLOGICAL ABSTRACTS INC. on STN Uridine 5'-diphospho-glucose-4-epimerase (UDP-Glc epimerase) catalyses the reversible epimerization of UDP-galactose and UDP-glucose. In contrast to bacteria and yeast, expression of the UDP-Glc epimerase gene in Arabidopsis was found not to be induced by galactose. To elucidate the metabolic role of this enzyme, transgenic Arabidopsis plants expressing the respective cDNA in sense or antisense orientation were constructed, leading to a range of plant lines with different UDP-Glc epimerase activities. No alterations in morphology were observed and the relative amounts of different galactose-containing compounds were not affected if the plants were raised on soil. However, on agar plates in the presence of galactose, the growth of different lines was increasingly repressed with decreasing enzyme activity, and an increase in the UDP-Gal content was observed in parallel, whereas the UDP-Glc content was nearly constant. The amount of galactose in the cell wall was increased in plants with low UDP-Glc epimerase activity grown on galactose, whereas the cellulose content in the leaves was not altered. Furthermore, starch determined at different times of the day was highly abundant in plants with low UDP-Glc epimerase activity in the presence of galactose. It is proposed that low endogenous UDP-Glc epimerase activity is responsible for the galactose toxicity of the wild-type. Possible mechanisms by which the starch content might be modulated are discussed.

L5 ANSWER 3 OF 3 CAPLUS COPYRIGHT 2003 ACS on STN Uridine 5'-diphospho-glucose-4-epimerase (UDP-Glc epimerase) catalyzes the AB reversible epimerization of UDP-galactose and UDP-glucose. In contrast to bacteria and yeast, expression of the UDP-Glc epimerase gene in Arabidopsis was found not to be induced by galactose. To elucidate the metabolic role of this enzyme, transgenic Arabidopsis plants expressing the resp. cDNA in sense or antisense orientation were constructed, leading to a range of plant lines with different UDP-Glc epimerase activities. No alterations in morphol. were obsd. and the relative amts. of different galactose-contg. compds. were not affected if the plants were raised on soil. However, on agar plates in the presence of galactose, the growth of different lines was increasingly repressed with decreasing enzyme activity, and an increase in the UDP Gal content was obsd. in parallel, whereas the UDP-Glc content was nearly const. The amt. of galactose in the cell wall was increased in plants with low UDP-Glc epimerase activity grown on galactose, whereas the cellulose content in the leaves was not

altered. Furthermore, starch detd. at different times of the day was highly abundant in plants with low UDP-Glc epimerase activity in the presence of galactose. It is proposed that low endogenous UDP-Glc epimerase activity is responsible for the galactose toxicity of the wild-type. Possible mechanisms by which the starch content might be modulated are discussed.

=> d 12 1-31

- ANSWER 1 OF 82 BIOSIS COPYRIGHT 2003 BIOLOGICAL ABSTRACTS INC. on STN DUPLICATE 1
- AN 2003:276389 BIOSIS
- DNPREV200300276389
- TTGALT deficiency causes UDP hexose deficit in human galactosemic cells.
- ΑU Lai, K. (1); Langley, S. D.; Khwaja, F. W.; Schmitt, E. W.; Elsas, L. J.
- CS (1) Department of Pediatrics, University of Miami School of Medicine, D-820, P.O. Box 016820, Miami, FL, 33101, USA: klai@med.miami.edu USA
- SO Glycobiology, (April 2003, 2003) Vol. 13, No. 4, pp. 285-294. print. ISSN: 0959-6658.
- DТ Article
- English LΑ
- L2 ANSWER 2 OF 82 BIOSIS COPYRIGHT 2003 BIOLOGICAL ABSTRACTS INC. on STN DUPLICATE 2
- AN 2001:231767 BIOSIS
- DN PREV200100231767
- Galactose metabolism in mice with galactose-1-phosphate uridyltransferase deficiency: Sucklings and 7-week-old animals fed a high-galactose diet.
- AH Ning, Cong; Reynolds, Robert; Chen, Jie; Yager, Claire; Berry, Gerard T.; Leslie, Nancy; Segal, Stanton (1)
- (1) Research Metabolism, Children's Hospital of Philadelphia, 3516 Civic Center Boulevard, 402 Abramson Pediatric Research Building, Philadelphia, PA, 19104-4318: segal@email.chop.edu USA
- Molecular Genetics and Metabolism, (April, 2001) Vol. 72, No. 4, pp. SO 306-315. print. ISSN: 1096-7192.
- DT Article
- English LA
- SLEnglish
- L2ANSWER 3 OF 82 CAPLUS COPYRIGHT 2003 ACS on STN
- AN2000:133844 CAPLUS
- DN 132:178178
- ΤI Galactose utilization as a positive selection marker in the transformation of plant cells
- ΙN Jorsboe, Morten; Brunstedt, Janne; Jorgensen, Kirsten
- PΑ Danisco A/S, Den.

- PCT Int. Appl., 86 pp. SO CODEN: PIXXD2
- DT Patent
- LA English
- FAN.CNT 1

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     Multiple copies of MRG19 suppress transcription of the GAL1 promoter in a
     GAL80-dependent manner in Saccharomyces cerevisiae.
ΑU
     Kabir, M. A.; Khanday, F. A.; Mehta, D. V.; Bhat, P. J. (1)
CS
     (1) Laboratory of Molecular Genetics, Biotechnology Center, Indian
     Institute of Technology, Powai Mumbai, 400 076 India
SO
     Molecular and General Genetics, (Jan., 2000) Vol. 262, No. 6, pp.
     1113-1122.
     ISSN: 0026-8925.
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L2
     ANSWER 5 OF 82 BIOSIS COPYRIGHT 2003 BIOLOGICAL ABSTRACTS INC. on STN
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     2001:54757 BIOSIS
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DN
     PREV200100054757
ΤI
     Studies of the V94M-substituted human UDPgalactose-4-epimerase enzyme
     associated with generalized epimerase-deficiency galactosaemia.
ΑU
     Wohlers, T. M.; Fridovich-Keil, J. L. (1)
CS
     (1) Department of Genetics, Emory University School of Medicine, 1462
     Clifton Rd, NE, Atlanta, GA, 30322: jfridov@emory.edu USA
     Journal of Inherited Metabolic Disease, (November, 2000) Vol. 23, No. 7,
SO
     pp. 713-729. print.
     ISSN: 0141-8955.
DT
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LA
     English
     English
SL
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     (2003) on STN
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     2000:71803 AGRICOLA
AN
DN
     IND22072400
     Expression of human inositol monophosphatase suppresses galactose
TI
     toxicity in Saccharomyces cerevisiae: possible implications in
     galactosemia.
     Mehta, D.V.; Kabir, A.; Bhat, P.J.
ΑU
     DNAL (381 B522)
ΑV
SO
     Biochimica et biophysica acta = International journal of biochemistry and
     biophysics, Aug 30, 1999. Vol. 1454, No. 3. p. 217-226
     Publisher: Amsterdam : Elsevier Science B.V.
     CODEN: BBACAQ; ISSN: 0006-3002
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DT
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FS
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LA
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- CS (1) Res. Inst., Hosp. Sick Children, 555 University Ave., Toronto, ON M5G 1X8 Canada

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- CS DIV. BIOCHEM., DEP. PHARM., JADAVPUR UNIV., CALCUTTA 700 032, INDIA.
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- CS DIV. PERINATAL MED., UNIV. COLORADO HEALTH SCI. CENT., CONTAINER B-199, 4200 E. NINTH AVE., DENVER, CO 80262.
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- TI THE EFFECT OF PHENTOLAMINE ON SYNAPTOSOMAL PHOSPHATIDYL INOSITOL IN EXPERIMENTAL GALACTOSE TOXICITY.
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- AU BERRY G; YANDRASITZ J R; SEGAL S
- CS CHILDREN'S HOSP. OF PHILADELPHIA, 34TH AND CIVIC CENTER BOULEVARD,

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- LA English
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- L6 0 GALACTOSE SELECTION AND MAMMAL
- => s galactose(w) selection and mammalian
- L7 0 GALACTOSE(W) SELECTION AND MAMMALIAN
- => s galactose(w)selection
- L8 10 GALACTOSE(W) SELECTION
- => d 18 1-10
- L8 ANSWER 1 OF 10 BIOSIS COPYRIGHT 2003 BIOLOGICAL ABSTRACTS INC. on STN
- AN 2003:281345 BIOSIS
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- TI A selection system for transgenic plants based on galactose as selective agent and a UDP-glucose:galactose-1-phosphate uridyltransferase gene as selective gene.
- AU Joersbo, Morten (1); Jorgensen, Kirsten; Brunstedt, Janne
- CS (1) Danisco Seed, Hojbygardvej 31, Holeby, DK-4960, Denmark: shmjädanisco.com Denmark
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- AU Miyazaki, Yoshitsugu; Geber, Antonia; Miyazaki, Haruko; Falconer, Derek; Parkinson, Tanya; Hitchcock, Christopher; Grimberg, Brian; Nyswaner, Katherine; Bennett, John E. (1)
- CS (1) Clinical Mycology Section, Laboratory of Clinical Investigation, National Institute of Allergy and Infectious Diseases, NIH, 10 Center Drive, Bethesda, MD, 20892 USA
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- LA English
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- AN 1986:142294 BIOSIS
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- AU WAGNER R P; COX S H; SCHOEN R C
- CS LIFE SCI. DIV., LS-3 GENETICS GROUP, MS M886, LOS ALAMOS NATIONAL LAB., LOS ALAMOS, NEW MEXICO 87545.
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- AU Miyazaki Y.; Geber A.; Miyazaki H.; Falconer D.; Parkinson T.; Hitchcock C.; Grimberg B.; Nyswaner K.; Bennett J.E.
- CS J.E. Bennett, Laboratory Clinical Investigation, National Inst. Allergy/Infect. Dis., NIH, 10 Center Drive, Bethesda, MD 20892, United States. jb46yanih.gov
- SO Gene, (1999) 236/1 (43-51). Refs: 32
 - ISSN: 0378-1119 CODEN: GENED6
- PUI S 0378-1119(99)00263-2
- CY Netherlands
- DT Journal; Article
- FS 004 Microbiology
- LA English
- SL English
- L8 ANSWER 5 OF 10 EMBASE COPYRIGHT 2003 ELSEVIER SCI. B.V. on STN
- AN 89171633 EMBASE
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- TI A novel genetic system to detect protein-protein interactions.
- AU Fields S.; Song O.-K.
- CS Department of Microbiology, State University of New York, Stony Brook, NY 11794, United States
- SO Nature, (1989) 340/6230 (245-246). ISSN: 0028-0836 CODEN: NATUAS
- CY United Kingdom
- DT Journal
- FS 004 Microbiology 029 Clinical Biochemistry
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- SL English
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- DN 1986064548

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      Chinese hamster.
      Wagner R.P.; Cox S.H.; Schoen R.C.
IJΔ
CS
      Life Sciences Division, LS-3, Genetics Group, Los Alamos National
      Laboratory, Los Alamos, NM 87545, United States
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ΑN
      2003:316379 CAPLUS
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     A selection system for transgenic plants based on galactose as selective
      agent and a UDP-glucose:galactose 1-phosphate uridyltransferase gene as
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ΑU
     Joersbo, Morten; Jorgensen, Kirsten; Brunstedt, Janne
     Danisco Seed, Holeby, DK-4960, Den.
CS
     Molecular Breeding (2003), 11(4), 315-323
     CODEN: MOBRFL; ISSN: 1380-3743
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     Kluwer Academic Publishers
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     Journal
     English
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     2002:675778 CAPLUS
DN
     137:213253
ΤI
     Selection by mirror image display
IN
     Wong, Chi-Huey
PΑ
     The Scripps Research Institute, USA
     PCT Int. Appl., 46 pp.
SO
     CODEN: PIXXD2
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     Patent
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     English
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     PATENT NO.
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     WO 2002067860
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         RW: GH, GM, KE, LS, MW, MZ, SD, SL, SZ, TZ, UG, ZM, ZW, AT, BE, CH,
              CY, DE, DK, ES, FI, FR, GB, GR, IE, IT, LU, MC, NL, PT, SE, TR,
              BF, BJ, CF, CG, CI, CM, GA, GN, GQ, GW, ML, MR, NE, SN, TD, TG
PRAI US 2001-271377P
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     1999:567893 CAPLUS
DN
     131:282268
     Cloning, sequencing, expression and allelic sequence diversity of ERG3
TI
     (C-5 sterol desaturase gene) in Candida albicans
ΑU
     Miyazaki, Yoshitsugu; Geber, Antonia; Miyazaki, Haruko; Falconer, Derek;
     Parkinson, Tanya; Hitchcock, Christopher; Grimberg, Brian; Nyswaner,
     Katherine; Bennett, John E.
     Clinical Mycology Section, Laboratory of Clinical Investigation, National
CS
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Institute of Allergy and Infectious Diseases, NIH, Bethesda, MD, 20892,

USA

SO Gene (1999), 236(1), 43-51 CODEN: GENED6; ISSN: 0378-1119

PB Elsevier Science B.V.

DT Journal

LA English

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AN 1989:530221 CAPLUS

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TI A novel genetic system to detect protein-protein interactions

AU Fields, Stanley; Song, Ok Kyu

CS Dep. Microbiol., State Univ. New York, Stony Brook, NY, 11794, USA

SO Nature (London, United Kingdom) (1989), 340 (6230), 245-6 CODEN: NATUAS; ISSN: 0028-0836

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L8 ANSWER 1 OF 10 BIOSIS COPYRIGHT 2003 BIOLOGICAL ABSTRACTS INC. on STN AB A new selection system based on galactose as selective agent and a

UDP-qlucose:galactose-1-phosphate uridyltransferase gene as selective gene is presented. A broad range of plant species, including agronomically important crops such as maize and rice, is sensitive to low dosages of galactose. The toxicity of galactose is believed to be due to accumulation of galactose-1-phosphate, generated by endogenous galactokinase after uptake. Here, it is demonstrated that this toxicity can be sufficiently alleviated by the Agrobacterium tumefaciens-mediated introduction of the E. coli UDP-glucose:galactose-1-phosphate uridyltransferase (galT) gene, driven by a 35S-promoter, to allow transgenic shoots of potato and oil seed rape to regenerate on galactose containing selection media, resulting in high transformation frequencies (up to 35% for potato). Analysis of genomic DNA and UDP-glucose:galactose-1-phosphate uridyltransferase activity in randomly selected potato transformants confirmed the presence and active expression of the galT gene. The agricultural performance of transgenic potatoes was evaluated by monitoring the phenotype and tuber yield for two generations and these characters were found to be indistinguishable from non-transgenic controls. Thus, the galactose selection system provides a new alternative being distinct from conventional antibiotic and herbicide selection systems as well as so-called positive selection systems where the selective agent has a beneficial effect.

ANSWER 2 OF 10 BIOSIS COPYRIGHT 2003 BIOLOGICAL ABSTRACTS INC. on STN AB The C-5 sterol desaturase gene (ERG3), essential for yeast ergosterol biosynthesis, was cloned and sequenced from Candida albicans by homology with the Saccharomyces cerevisiae ERG3. The ERG3 ORF contained 1158 bp and encoded 386 deduced amino acids. The clone was used to transform a gall mutant derived from the Darlington strain of C. albicans, using galactose selection. The Darlington strain is known to lack DELTA5,6 sterols, i.e. to have an erg3 phenotype (Howell, S.A., et al., 1990. J. Appl. Bacteriol. 69, 692-696). The transformant (CDTR1) contained six tandem integrated ERG3GAL1 repeats, had double the abundance of ERG3 transcript found in the host strain, and synthesized ergosterol, a DELTA5,6 sterol. The Darlington strain was noted to have an abundance of ERG3 transcript. Both ERG3 alleles in Darlington were cloned and sequenced in order to look for changes that might explain the erg3 phenotype. One allele, called Dar-2, contained a stop codon in place of tryptophan-292. The other ERG3 allele, called Dar-1, had changes in three amino acids, two of which were conserved in three fungal and one plant species. EcoRI genomic fragments containing ERG3 from the Dar-1 allele and from B311, the wild type strain, were inserted into the plasmid pRS316 and used to transform a Saccharomyces cerevisiae erg3, ura3 mutant using uracil selection. The 4.1 kb ERG3 fragments from the B311 and Dar-1 both contained 1.4 kb 5' and 1.5 kb 3' flanking sequences around the coding region. Transformants with ERG3 from B311 but not from Dar-1 showed restored ergosterol synthesis. One or more of these three deduced amino acids in the Dar-1 allele of ERG3 appeared critical for function.

ANSWER 3 OF 10 BIOSIS COPYRIGHT 2003 BIOLOGICAL ABSTRACTS INC. on STN Chinese hamster cells in culture were treated with various concentrations of thymidine, 5-bromodeoxyuridine, trifluorothymidine, and 2-deoxy-Dgalactose. Selection was made for deficiencies in the activities of galactokinase and thymidine kinase. Selection in the presence of thymidine, 5-bromodeoxyuridine, and trifluorothymidine was expected to produce clones deficient in thymidine kinase only, whereas those deficient in galactokinase were expected to be selected in the presence of 2-deoxy-D-galactose. However, it was found that clones growing in the presence of these inhibitors were frequently deficient in both enzymes. Or if a clone was deficient in only one, the deficiency frequently was not expected according to the selection procedure. This indicates some sort of coordinate relationship between the two gene loci, GALK and TK1, which specify galactokinase and thymidine kinase, respectively. GALK and TK1 are linked in all primates and rodents in which linkage determinations have been made. It is therefore probable that this

L8

linkage has been conserved for a long period of time. It is suggested that the apparent relationship between the two genes shown by the data presented here, as well as by others, supports the conclusion that linkage has been conserved by natural selection and is therefore not fortuitous.

- ANSWER 4 OF 10 EMBASE COPYRIGHT 2003 ELSEVIER SCI. B.V. on STN 1.8 The C-5 sterol desaturase gene (ERG3), essential for yeast ergosterol AB biosynthesis, was cloned and sequenced from Candida albicans by homology with the Saccharomyces cerevisiae ERG3. The ERG3 ORF contained 1158 bp and encoded 386 deduced amino acids. The clone was used to transform a gall mutant derived from the Darlington strain of C. albicans, using galactose selection. The Darlington strain is known to lack .DELTA.5,6 sterols, i.e. to have an erg3 phenotype (Howell, S.A., et al., 1990. J. Appl. Bacteriol. 69, 692-696). The transformant (CDTR1) contained six tandem integrated ERG3GAL1 repeats, had double the abundance of ERG3 transcript found in the host strain, and synthesized ergosterol, a .DELTA.5,6 sterol. The Darlington strain was noted to have an abundance of ERG3 transcript. Both ERG3 alleles in Darlington were cloned and sequenced in order to look for changes that might explain the erg3 phenotype. One allele, called Dar-2, contained a stop codon in place of tryptophan-292. The other ERG3 allele, called Dar-1, had changes in three amino acids, two of which were conserved in three fungal and one plant species. EcoRI genomic fragments containing ERG3 from the Dar-1 allele and from B311, the wild-type strain, were inserted into the plasmid pRS316 and used to transform a Saccharomyces cerevisiae erg3, ura3 mutant using uracil selection. The 4.1 kb ERG3 fragments from the B311 and Dar-1 both contained 1.4 kb 5' and 1.5 kb 3' flanking sequences around the coding region. Transformants with ERG3 from B311 but not from Dar-1 showed restored ergosterol synthesis. One or more of these three deduced amino acids in the Dar-1 allele of ERG3 appeared critical for function. (C) 1999 Published by Elsevier Science B.V. All rights reserved.
- ANSWER 5 OF 10 EMBASE COPYRIGHT 2003 ELSEVIER SCI. B.V. on STN Protein-protein interactions between two proteins have generally been ΔR studied using biochemical techniques such as crosslinking, co-immunoprecipitation and co-fractionation by chromatography. We have generated a novel genetic system to study these interactions by taking advantage of the properties of the GAL4 protein of the yeast Saccharomyces cerevisiae. This protein is a transcriptional activator required for the expression of genes encoding enzymes of galactose utilization. It consists of two separable and functionally essential domains: an N-terminal domain which binds to specific DNA sequences (UAS(G)); and a C-terminal domain containing acidic regions, which is necessary to activate transcription. We have generated a system of two hybrid proteins containing parts of GAL4: the GAL4 DNA-binding domain fused to a protein 'X' and a GAL4 activating region fused to a protein 'Y'. If X and Y can form a protein-protein complex and reconstitute proximity of the GAL4 domains, transcription of a gene regulated by UAS(G) occurs. We have tested this system using two yeast proteins that are known to interact-SNF1 and SNF4. High transcriptional activity is obtained only when both hybrids are present in a cell. This system may be applicable as a general method to identify proteins that interact with a known protein by the use of a simple galactose selection.
- ANSWER 6 OF 10 EMBASE COPYRIGHT 2003 ELSEVIER SCI. B.V. on STN

 Chinese hamster cells in culture were treated with various concentrations of thymidine, 5-bromodeoxyuridine, trifluorothymidine, and 2-deoxy-D-galactose. Selection was made for deficiencies in the activities of galactokinase and thymidine kinase. Selection in the presence of thymidine, 5-bromodeoxyuridine, and trifluorothymidine was expected to produce clones deficient in thymidine kinase only, whereas those deficient in galactokinase were expected to be selected in the presence of 2-deoxy D-galactose. However, it was found that clones growing in the presence of these inhibitors were frequently deficient in both

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- ANSWER 7 OF 10 CAPLUS COPYRIGHT 2003 ACS on STN 1.8 AΒ A new selection system based on galactose as selective agent and a UDP-qlucose:galactose-1-phosphate uridyltransferase gene as selective gene is presented. A broad range of plant species, including agronomically important crops such as maize and rice, is sensitive to low dosages of galactose. The toxicity of galactose is believed to be due to accumulation of galactose-1-phosphate, generated by endogenous galactokinase after uptake. Here, it is demonstrated that this toxicity can be sufficiently alleviated by the Agrobacterium tumefaciens-mediated introduction of the E. coli UDP-glucose:galactose-1-phosphate uridyltransferase (galT) gene, driven by a 35S-promoter, to allow transgenic shoots of potato and oil seed rape to regenerate on galactose contg. selection media, resulting in high transformation frequencies (up to 35% for potato). Anal. of genomic DNA and UDP-qlucose:galactose-1phosphate uridyltransferase activity in randomly selected potato transformants confirmed the presence and active expression of the galT gene. The agricultural performance of transgenic potatoes was evaluated by monitoring the phenotype and tuber yield for two generations and these characters were found to be indistinguishable from non-transgenic controls. Thus, the galactose selection system provides a new alternative being distinct from conventional antibiotic and herbicide selection systems as well as so-called pos. selection systems where the selective agent has a beneficial effect.
- ANSWER 8 OF 10 CAPLUS COPYRIGHT 2003 ACS on STN

 Non-naturally occurring binders to cell surface carbohydrates and sugars are identified by a screening process that employs immobilized enantiomers of such cell surface carbohydrates and sugars. Preferred non-naturally occurring binders include D-peptides and L-nucleic acids and are resistant to enzymic degrdn. and clearance. Single-chain Fab sequences that bind to sialic acid and KDO in nano-molar affinity were identified by this process. Exemplary screening procedures employed D-KDO, L-sialic acid and an L-sialo-disaccharide have been attached to a solid support for selection of high-affinity binders.
- 1.8 ANSWER 9 OF 10 CAPLUS COPYRIGHT 2003 ACS on STN The C-5 sterol desaturase gene (ERG3), essential for yeast ergosterol AB biosynthesis, was cloned and sequenced from Candida albicans by homol. with the Saccharomyces cerevisiae ERG3. The ERG3 ORF contained 1158 bp and encoded 386 deduced amino acids. The clone was used to transform a gall mutant derived from the Darlington strain of C. albicans, using galactose selection. The Darlington strain is known to lack .DELTA.5,6 sterols, i.e. to have an erg3 phenotype (Howell, S.A., et al., 1990. J. Appl. Bacteriol. 69, 692-696). The transformant (CDTR1) contained six tandem integrated ERG3GAL1 repeats, had double the abundance of ERG3 transcript found in the host strain, and synthesized ergosterol, a .DELTA.5,6 sterol. The Darlington strain was noted to have an abundance of ERG3 transcript. Both ERG3 alleles in Darlington were cloned and sequenced in order to look for changes that might explain the erg3 phenotype. One allele, called Dar-2, contained a stop codon in place of tryptophan-292. The other ERG3 allele, called Dar-1, had changes in three amino acids, two of which were conserved in three fungal and one plant species. EcoRI genomic fragments contg. ERG3 from the Dar-1 allele and

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